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Asp Asn Val Lys Pro Ile Tyr Ile Val Thr Pro Thr Asn Ala Ser His 50 55 60

Ile Gln Ser Ala Val Val Cys Gly Arg Arg His Gly Val Arg Ile Arg 65 70 75 80

Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu Ser Tyr Arg Ser Leu 85 90 95

Gln Pro Glu Glu Phe Ala Val Val Asp Leu Ser Lys Met Arg Ala Val
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Trp Val Asp Gly Lys Ala Arg Thr Ala Trp Val Asp Ser Gly Ala Gln
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Leu Gly Glu Leu Tyr Tyr Ala Ile His Lys Ala Ser Thr Val Leu Ala 130 135 140

Phe Pro Ala Gly Val Cys Pro Thr Ile Gly Val Gly Gly Asn Phe Ala 145 150 155 160

Gly Gly Phe Gly Met Leu Leu Arg Lys Tyr Gly Ile Ala Ala Glu 165 170 175

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Gly Glu Ser Phe Gly Ile Val Val Ala Trp Lys Val Arg Leu Leu Pro 210 215 220

Val Pro Pro Thr Val Thr Val Phe Lys Ile Pro Lys Lys Ala Ser Glu 225 230 235 240

Gly Ala Val Asp Ile Ile Asn Arg Trp Gln Val Val Ala Pro Gln Leu 245 250 255

Pro Asp Asp Leu Met Ile Arg Val Ile Ala Gln Gly Pro Thr Ala Thr 260 265 270

Phe Glu Ala Met Tyr Leu Gly Thr Cys Gln Thr Leu Thr Pro Met Met 275 280 285

Ser Ser Lys Phe Pro Glu Leu Gly Met Asn Ala Ser His Cys Asn Glu 290 295 300

Met Ser Trp Ile Gln Ser Ile Pro Phe Val His Leu Gly His Arg Asp 305 310 315 320

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Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Glu Pro Phe Pro Lys Arg Val 340 345 350

Trp Glu Gln Ile Phe Ser Thr Trp Leu Leu Lys Pro Gly Ala Gly Ile 355 360 365

Met Ile Phe Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro Glu Trp 370 375 380

Ala Thr Pro Phe Pro His Arg Lys Gly Val Leu Phe Asn Ile Gln Tyr 385 390 395 400

Val Asn Tyr Trp Phe Ala Pro Gly Ala Gly Ala Ala Pro Leu Ser Trp 405 410 415

Ser Lys Glu Ile Tyr Asn Tyr Met Glu Pro Tyr Val Ser Lys Asn Pro 420 425 430

A. .

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					tac Tyr	_										432
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	Pro				acc Thr 230											720
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ttc	gag	gcc	atg	tac	ctg	ggc	acc	tgc	caa	acc	ctg	acg	ccg	atg	atg	864

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Asp Asn Val Lys Pro Ile Tyr Ile Val Thr Pro Thr Asn Ala Ser His 50 55 60

Ile Gln Ser Ala Val Val Cys Gly Arg Arg His Gly Val Arg Ile Arg 65 70 75 80

Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu Ser Tyr Arg Ser Leu 85 90 95

Gln Pro Glu Glu Phe Ala Val Val Asp Leu Ser Lys Met Arg Ala Val 100 105 110

Trp Val Asp Gly Lys Ala Arg Thr Ala Trp Val Asp Ser Gly Ala Gln
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Phe Pro Ala Gly Val Cys Pro Thr Ile Gly Val Gly Gly Asn Phe Ala 145 150 155 160

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Asn Val Ile Asp Val Lys Leu Val Asp Ala Asn Gly Thr Leu His Asp 180 185 190 Lys Lys Ser Met Gly Asp Asp His Phe Trp Ala Val Arg Gly Gly Gly 195 200 205

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Val Pro Pro Thr Val Thr Val Phe Lys Ile Pro Lys Lys Ala Ser Glu 225 230 235 240

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Pro Asp Asp Leu Met Ile Arg Val Ile Ala Gln Gly Pro Thr Ala Thr 260 265 270

Phe Glu Ala Met Tyr Leu Gly Thr Cys Gln Thr Leu Thr Pro Met Met 275 280 285

Ser Ser Lys Phe Pro Glu Leu Gly Met Asn Ala Ser His Cys Asn Glu 290 295 300

Met Ser Trp Ile Gln Ser Ile Pro Phe Val His Leu Gly His Arg Asp 305 310 315 320

Asn Ile Glu Asp Asp Leu Leu Asn Arg Asn Asn Thr Phe Lys Pro Phe 325 330 335

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Trp Glu Gln Ile Phe Ser Thr Trp Leu Leu Lys Pro Gly Ala Gly Ile 355 360 365

Met Ile Phe Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro Glu Trp 370 375 380

Ala Thr Pro Phe Pro His Arg Lys Gly Val Leu Phe Asn Ile Gln Tyr 385 390 395 400

Val Asn Tyr Trp Phe Ala Pro Gly Ala Gly Ala Ala Pro Leu Ser Trp 405 410 415 Ser Lys Glu Ile Tyr Asn Tyr Met Glu Pro Tyr Val Ser Lys Asn Pro 430 425 420 Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly Arg Asn Glu 440 435 Val Val Asn Asp Val Ser Thr Phe Ser Ser Gly Leu Val Trp Gly Gln 450 455 460 Lys Tyr Phe Lys Gly Asn Phe Gln Arg Leu Ala Ile Thr Lys Gly Lys 470 475 480 Val Asp Pro Thr Asp Tyr Phe Arg Asn Glu Gln Ser Ile Pro Pro Leu 490 Ile Lys Lys Tyr 500 <210> 5 <211> 1503 <212> DNA <213> Phleum pratense <220> <221> artificial_DNA_sequence <222> (1)..(69) <223> DNA sequence derived from sequenced protein <220> <221> native_DNA_sequence <222> (70)..(1503) <223> <220> <221> CDS <222> (1)..(1503) <223> tac ttc ccg ccg ccg gct gct aaa gac ttc ctg ggt tgc ctg gtt 48 Tyr Phe Pro Pro Pro Ala Ala Lys Glu Asp Phe Leu Gly Cys Leu Val aaa gaa atc ccg ccg cgt ctg ttg tac gcg aaa tcg tcg ccg gcg tat 96 Lys Glu Ile Pro Pro Arg Leu Leu Tyr Ala Lys Ser Ser Pro Ala Tyr 144 ccc tca gtc ctg ggg cag acc atc cgg aac tcg agg tgg tcg tcg ccg

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						ata Ile										720
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Asp Asn Val Lys Pro Leu Tyr Ile Ile Thr Pro Thr Asn Val Ser His 50 55 60

Ile Gln Ser Ala Val Val Cys Gly Arg Arg His Ser Val Arg Ile Arg 65 70 75 80

Val Arg Ser Gly Gly His Asp Tyr Glu Gly Leu Ser Tyr Arg Ser Leu 85 90 95

Gln Pro Glu Thr Phe Ala Val Val Asp Leu Asn Lys Met Arg Ala Val 100 105 110

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Lys Lys Ser Met Gly Asp Asp His Phe Trp Ala Val Arg Gly Gly 195 200 205

Gly Glu Ser Phe Gly Ile Val Val Ala Trp Gln Val Lys Leu Leu Pro 210 215 220

Val Pro Pro Thr Val Thr Ile Phe Lys Ile Ser Lys Thr Val Ser Glu 225 230 235 240

Gly Ala Val Asp Ile Ile Asn Lys Trp Gln Val Val Ala Pro Gln Leu 245 250 255

Pro Ala Asp Leu Met Ile Arg Ile Ile Ala Gln Gly Pro Lys Ala Thr 260 265 270

Phe Glu Ala Met Tyr Leu Gly Thr Cys Lys Thr Leu Thr Pro Leu Met 275 280 285

Ser Ser Lys Phe Pro Glu Leu Gly Met Asn Pro Ser His Cys Asn Glu 290 295 300

Met Ser Trp Ile Gln Ser Ile Pro Phe Val His Leu Gly His Arg Asp 305 310 315 320

Ala Leu Glu Asp Asp Leu Leu Asn Arg Asn Asn Ser Phe Lys Pro Phe 325 330 335

Ala Glu Tyr Lys Ser Asp Tyr Val Tyr Gln Pro Phe Pro Lys Thr Val 340 345 350

Trp Glu Gln Ile Leu Asn Thr Trp Leu Val Lys Pro Gly Ala Gly Ile 355 360 365

Met Ile Phe Asp Pro Tyr Gly Ala Thr Ile Ser Ala Thr Pro Glu Ser 370 375 380

Ala Thr Pro Phe Pro His Arg Lys Gly Val Leu Phe Asn Ile Gln Tyr 385 390 395 400

Val Asn Tyr Trp Phe Ala Pro Gly Ala Ala Ala Pro Leu Ser Trp

405 410 415

Ser Lys Asp Ile Tyr Asn Tyr Met Glu Pro Tyr Val Ser Lys Asn Pro 420 425

Arg Gln Ala Tyr Ala Asn Tyr Arg Asp Ile Asp Leu Gly Arg Asn Glu 440

Val Val Asn Asp Val Ser Thr Tyr Ala Ser Gly Lys Val Trp Gly Gln 455

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See.

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